

Table 4. Selected *ASPM* Normal Variants

DNA Nucleotide Change (Alias ¹)	Protein Amino Acid Change (Alias ¹)	Reference Sequences
c.848C>T	p.= (p.Ser283Ser)	
c.937A>G	p.Ile313Val	
c.1288A>G	p.Arg430Gly	
c.1977T>C	p.= (p.Ile659Ile)	
c.2606C>T	p.Thr869Ile	
c.2779T>C	p.= (p.Leu927Leu)	
c.2805T>C	p.= (p.Ser935Ser)	
c.3138 G>A	p.= (p.Arg1046Arg)	
c.3269C>T	p.Ser1090Phe	
c.3579T>A	p.= p.Ser1193Ser	
c.4449A>G	p.= (p.Lys1483Lys)	NM_018136.4 NP_060606.3
c.5961A>G	p.Gln2147Gln	
c.7461T>A	p.= (p.Ile2487Ile)	
c.7480T>C	p.Tyr2494His	
c.7566A>G	p.= (p. Leu2522Leu)	
c.7605G>A	p.= (p.Val2535Val)	
c.7674 C>T	p.= (p.Ile2558Ile)	
c.7684A>G	p.Ser2562Gly	
c.7860G>C	p.Gln2620His	
c.7939C>A	p.Leu2647Ile	
c.9395T>G	p.Leu3132Arg	
c.9773A>G	p.His3258Arg	

See [Quick Reference](#) for an explanation of nomenclature. *GeneReviews* follows the standard naming conventions of the Human Genome Variation Society (<http://www.hgvs.org>).

1. Variant designation that does not conform to current naming conventions